

**Amendment to the Claims**

The following listing of claims will replace all prior versions and listings of claims.

**Listing of Claims:**

1. (Currently Amended) An isolated nucleic acid molecule comprising a first polynucleotide sequence at least 15% identical to a second polynucleotide sequence selected from the group consisting of:
  - (a) a polynucleotide fragment of SEQ ID NO:132 as referenced in Table 1A;
  - (b) a polynucleotide encoding a full length polypeptide of SEQ ID NO:421, or a full length polypeptide encoded by the HODFN71 cDNA Clone ID in ATCC Deposit No:203570, corresponding to SEQ ID NO:421 as referenced in Table 1A;
  - (c) a polynucleotide encoding a polypeptide fragment of SEQ ID NO:421 or a polypeptide fragment encoded by the HODFN71 cDNA Clone ID in ATCC Deposit No:203570 corresponding to SEQ ID NO:421 as referenced in Table 1A;
  - (d) a polynucleotide encoding a polypeptide fragment of SEQ ID NO:421 or a polypeptide fragment encoded by the HODFN71 cDNA Clone ID in ATCC Deposit No:203570 corresponding to SEQ ID NO:421 as referenced in Table 1A, wherein said fragment has biological activity;
  - (e) a polynucleotide encoding a predicted epitope of SEQ ID NO:421 as referenced in Table 1B; and
  - (f) a polynucleotide capable of hybridizing under stringent conditions to any one of the polynucleotides specified in (a) - (e), wherein said polynucleotide does not hybridize under stringent conditions to a nucleic acid molecule having a nucleotide sequence of only A residues or of only T residues.

2 - 24. (Cancelled)

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